RAW SEQUENCE LISTING DATE: 04/18/2003
PATENT APPLICATION: **US/09/972,758A** TIME: 15:54:26

Input Set : **A:\277084004.ST25.txt**
Output Set: **N:\CRF4\04182003\I972758A.raw**

<110> APPLICANT: Case Western Reserve University
Montano, Monica
Wittman, Bryan
<120> TITLE OF INVENTION: Suppressors of Human Breast Cancer Cell Growth
<130> FILE REFERENCE: 27708/04004
<140> CURRENT APPLICATION NUMBER: US 09/972758A
<141> CURRENT FILING DATE: 2001-10-05
<150> PRIOR APPLICATION NUMBER: US 60/238,187
<151> PRIOR FILING DATE: 2000-10-05
<160> NUMBER OF SEQ ID NOS: 7
<170> SOFTWARE: PatentIn version 3.1

<210> SEQ ID NO: 1
<211> LENGTH: 1080
<212> TYPE: DNA
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 1

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cccgaggagg acagtaggtg gcaatcgaga gcgttccccc agttgggtgg ccgtccgggg    180
ccggaggggg aagggagcct ggaatcccaa ccacctccct tgcagaccca ggcctgtcca    240
gaatctagct gcctgagaga gggcgagaag ggccagaatg gggacgactc gtccgctggc    300
ggcgacttcc cgccgccggc agaagtggaa ccgacgcccg aggccgagct gctcgcccag    360
ccttgtcatg actccgaggc cagtaagttg ggggctcctg ccgcaggggg cgaagaggag    420
tggggacagc agcagagaca gctggggaag aaaaaacata agagacgccc gtccaagaag    480
aagcggcatt ggaaaccgta ctacaagctg aactgggaag agaagaaaaa gttcgacgag    540
aaacagagcc ttcgagcttc aaggatccga gccgagatgt tcgccaaggg ccagccggtc    600
gcgccctata acaccacgca gttcctcatg gatgatcacg accaggagga gccggatctc    660
aaaaccggcc tgtactccaa gcgggccgcc gccaaatccg acgacaccag cgatgacgac    720
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gagtttctgc agcgggactt ctcggagacg tacgagcggt accacacgga gagcctgcag    840
aacatgagca agcaggagct catcaaggag tacctggaac tggagaagtg cctctcgcgc    900
atggaggacg agaacaaccg gctgcggctg gagagcaagc ggctgggtgg cgacgacgcg    960
cgtgtgcggg agctggagct ggagctggac cggctgcgcg ccgagaacct ccagctgctg   1020
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<210> SEQ ID NO: 2
<211> LENGTH: 359
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 2

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Met Ala Glu Pro Phe Leu Ser Glu Tyr Gln His Gln Pro Gln Thr Ser
1               5                  10                  15
Asn Cys Thr Gly Ala Ala Ala Val Gln Glu Glu Leu Asn Pro Glu Arg
            20                  25                  30
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 79 Pro Pro Gly Ala Glu Glu Arg Val Pro Glu Glu Asp Ser Arg Trp Gln
 80         35                  40                  45
 83 Ser Arg Ala Phe Pro Gln Leu Gly Gly Arg Pro Gly Pro Glu Gly Glu
 84     50                  55                  60
 87 Gly Ser Leu Glu Ser Gln Pro Pro Pro Leu Gln Thr Gln Ala Cys Pro
 88 65                  70                  75                  80
 91 Glu Ser Ser Cys Leu Arg Glu Gly Glu Lys Gly Gln Asn Gly Asp Asp
 92                 85                  90                  95
 95 Ser Ser Ala Gly Gly Asp Phe Pro Pro Pro Ala Glu Val Glu Pro Thr
 96             100                 105                 110
 99 Pro Glu Ala Glu Leu Leu Ala Gln Pro Cys His Asp Ser Glu Ala Ser
100         115                 120                 125
103 Lys Leu Gly Ala Pro Ala Ala Gly Gly Glu Glu Glu Trp Gly Gln Gln
104     130                 135                 140
107 Gln Arg Gln Leu Gly Lys Lys Lys His Arg Arg Arg Pro Ser Lys Lys
108 145                 150                 155                 160
111 Lys Arg His Trp Lys Pro Tyr Tyr Lys Leu Thr Trp Glu Glu Lys Lys
112                 165                 170                 175
115 Lys Phe Asp Glu Lys Gln Ser Leu Arg Ala Ser Arg Ile Arg Ala Glu
116             180                 185                 190
119 Met Phe Ala Lys Gly Gln Pro Val Ala Pro Tyr Asn Thr Thr Gln Phe
120         195                 200                 205
123 Leu Met Asp Asp His Asp Gln Glu Glu Pro Asp Leu Lys Thr Gly Leu
124     210                 215                 220
127 Tyr Ser Lys Arg Ala Ala Ala Lys Ser Asp Asp Thr Ser Asp Asp Asp
128 225                 230                 235                 240
131 Phe Met Glu Glu Gly Gly Glu Glu Asp Gly Gly Ser Asp Gly Met Gly
132                 245                 250                 255
135 Gly Asp Gly Ser Glu Phe Leu Gln Arg Asp Phe Ser Glu Thr Tyr Glu
136             260                 265                 270
139 Arg Tyr His Thr Glu Ser Leu Gln Asn Met Ser Lys Gln Glu Leu Ile
140         275                 280                 285
143 Lys Glu Tyr Leu Glu Leu Glu Lys Cys Leu Ser Arg Met Glu Asp Glu
144     290                 295                 300
147 Asn Asn Arg Leu Arg Leu Glu Ser Lys Arg Leu Gly Gly Asp Asp Ala
148 305                 310                 315                 320
151 Arg Val Arg Glu Leu Glu Leu Glu Leu Asp Arg Leu Arg Ala Glu Asn
152                 325                 330                 335
155 Leu Gln Leu Leu Thr Glu Asn Glu Leu His Arg Gln Gln Glu Arg Ala
156             340                 345                 350
159 Pro Leu Ser Lys Phe Gly Asp
160         355
163 <210> SEQ ID NO: 3
164 <211> LENGTH: 19
165 <212> TYPE: PRT
166 <213> ORGANISM: Homo sapiens
168 <400> SEQUENCE: 3
170 Lys His Arg Arg Arg Pro Ser Lys Lys Lys Arg His Trp Lys Pro Tyr
171 1               5                   10                  15
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174 Tyr Lys Leu
178 <210> SEQ ID NO: 4
179 <211> LENGTH: 18
180 <212> TYPE: DNA
181 <213> ORGANISM: Homo sapiens
183 <400> SEQUENCE: 4
184 cagtgtgatt tctagagc 18
187 <210> SEQ ID NO: 5
188 <211> LENGTH: 18
189 <212> TYPE: DNA
190 <213> ORGANISM: Homo sapiens
192 <400> SEQUENCE: 5
193 agagcagaac tactcaag 18
196 <210> SEQ ID NO: 6
197 <211> LENGTH: 38
198 <212> TYPE: DNA
199 <213> ORGANISM: Homo sapiens
201 <400> SEQUENCE: 6
202 acacaggatc cgaattcatg tccggagccc ttgatgtc 38
205 <210> SEQ ID NO: 7
206 <211> LENGTH: 42
207 <212> TYPE: DNA
208 <213> ORGANISM: Homo sapiens
210 <400> SEQUENCE: 7
211 acacaggatc cagtcgacta agaccagtca gtggttgctc ct 42

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